



Figure 3

	SEQ ID NO:	Sequence Type	Binding Site
NLVATCLPVRASLPHRLN	[SEQ ID NO: 13]	5' untranslated region	
1MLRGPGLLLAVQCLGTAVPSTGASKSR	[SEQ ID NO: 14]	Signal	
32QAQQWVQPSVAVSQSKPG	[SEQ ID NO: 15]	NH2-terminal segment	
52CYDNGKHYYQINQWERTYLGNALVCTCYGSGRGFNCESKPEAEET	[SEQ ID NO: 16]	I	Fibrin Heparin S. aureus
97CFDKYTGNTYRVGDTYERPKDSMIMDCTCIGAGRGRI SCTIANR	[SEQ ID NO: 38]	I	
141CHEGGQSYKIGDTWRARPHETGGYMLECVCLGNGKGEMTCKPIAEK	[SEQ ID NO: 39]	I	
186CFDHAAGTSYVVGETWEKPYQGMMVDCCTCLGEGSGRITGTSRNR	[SEQ ID NO: 40]	I	
232CNDQDTRTSYRIGDTWSKKDNRGNLQICICTGNGRGEMKCEK	[SEQ ID NO: 17]	I	
273HTSVQTTSSGSGPFTDVRAVYQPDHPQPPYGH	[SEQ ID NO: 18]	Connecting strand	
303CVTDSGVVYVSGMQMLKTQGNKQMLCTCLGNGVSCQE	[SEQ ID NO: 19]	I	
343TAVTQTYGNSNGEPCVLFTYNDRTDSTSNYEQDQKYSFCTDH	[SEQ ID NO: 20]	II	Gelatin
390TVLVQTRGNSNGALCHFPFLYNNHYTDC TSEGRDNNKMGCTT	[SEQ ID NO: 28]	II	
QNYDADQKFGFCPMAHEEI			
453CTTNEGVMYRIGDQMDKQDHMGHMRCTCVGNGRGEMTCTIAYSQLRDQ	[SEQ ID NO: 21]	I	
503CIYDDITYNVNDTFHHRHEEGHMLNCTCFGGGRGWKCDPVDQ	[SEQ ID NO: 22]	I	
543CQDSETGT FYQIGDSWEKYYHGVRYQCYGGRGIGEMHCQPLQTYPSS	[SEQ ID NO: 23]	I	
594SGEVEVFITETPSQPNSHPIQWNAFPQPSHISKYILRWRP	[SEQ ID NO: 24]	III	
633VSIIPRNLGY	[SEQ ID NO: 25]	Unique Sequence	

*VS*SYQF*WLFEPAPFPTLLINYSYSIYYICLVNKQYVNV*IDL*TEKKKKKK[SEQ ID NO: 29-33]3' untranslated region